61

CCCGGAGCTAAGGCGCCGAACCCGCGGCGGCGGTGGGGACGATGTGGTTTTTTGCCCGG

GACCCGGTCCGGGACTTTCCGTTCGAGCTCATCCCGGAGCCCCCAGAGGGCGGCCTGCCC

121 GGGCCCTGGGCCCTGCACCGCGCCGCAAGAAGGCCACAGGCAGCCCCGTGTCCATCTTC 181 GTCTATGATGTGAAGCCTGCGCGGAARGAGCAGACCCAGGTGGCCAAAGCTGCCTTCAA 241 GCRGCTTCAAAACTCTACGGCACCCCAACATCRCTGGCTTACATCGATGGACTGGAGACA GAAAAATGCCTCACGTCGTGACAGAGGCTGTGACCCCGTTGGGAATATACCTCAAGGCG 301 361 AGAGTGGAGGCTGGTGGCCTGAAGGAGCTGGAGATCTCCTGGGGGCTACACCAGATCGTG 421 AAAGCCCTCAGCTTCCTGGTCAACGACTGCAGCCTCATCCACAACAATGTCTGCATGGCC 481 GCCGTGTTCGTGGACCGAGCTGGCGAGTGGAAGCTTGGGGGCCTGGACTACATGTATTCG GCCCAGGGCAACGGTGGGGGACCTCCCCGCAAGGGGATCCCCGAGCTTGAGCAGTATGAC 541 601 661 CTACGCAACCCTGGGAAGATCCCCAAAACGCTGGCGCCCCATTACTGTGAGCTGGTGGGA 721 GCAAACCCCAAGGTGCGTCCCAACCCAGCCCGCTTCCTGCAGAACTGCCGGGCACCTGGT 781 GGCTTCATGAGCAACCGCTTTGTAGAAACCAACCTCTTCCTGGAGGAGATTCAGATCAAA 841 GAGCCAGCCGAGAAGCAAAAATTCTTCCAGGAGCTGAGCAAGAGCCTGGACGCATTCCCT 901 GAGGATTTCTGTCGGCACAAGCTGCTGCCCCAGCTGCTGACCGCCTTCGAGTTCGGCAAT 961 1021 GCTGGGGCCGTTGTCCTCACGCCCCTCTTCAAGGTGGGCAAGTTCCTGAGCGCTGAGGAG 1081 TATCAGCAGAAGATCATCCCTGTGGTGGTCAAGATGTTCTCATCCACTGACCGGGCCATG 1141 CGCATCCGCCTCCTGCAGCAGATGGAGCAGTTCATCCAGTACCTTGACGAGCCAACAGTC 1201 AACACCCAGATCTTCCCCCACGTCGTACATGGCTTCCTGGACACCCAACCCTGCCATCCGG 1261 GAGCAGACGGTCAAGTCCATGCTGCTCCTGGCCCCAAAGCTGAACGAGGCCAACCTCAAT 1321 GTGGAGCTGATGAAGCACTTTGCACGGCTACAGGCCAAGGATGAACAGGGCCCCATCCGC 1381 TGCAACACCACAGTCTGCCTGGGCAAAATCGGCTCCTACCTCAGTGCTAGCACCAGACAC 1441 AGGGTCCTTACCTCTGCCTTCAGCCGAGCCACTAGGGACCCGTTTGCACCGTCCCGGGTT 1501 GCGGGTGTCCTGGGCTTTGCTGCCACCCACACCTCTACTCAATGAACGACTGTGCCCAG 1561 AAGATCCTGCCTGTGCTCTGCGGTCTCACTGTAGATCCTGAGAAATCCGTGCGAGACCAG 1621 GCCTTCAAGGCATTTCGGAGCTTCCTGTCCAAATTGGAGTCTGTGTCGGAGGACCCGACC 1681 CAGCTGGAGGAAGTGGAGAAGGATGTCCATGCAGCCTCCAGCCCTGGCATGGGAGGAGCC

FIGURE 1A

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FIGURE 1B

1 .	O.G. FIG.	
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NSGNNAEEAPGAKAPEPAAAVGTMWFFARDPVRDFPFELIPEPPEGGLPGPWALHRGRKK 1 ATGSPVSIFVYDVKPGAEEQTQVAKAAFKRFKTLRHPNILAYIDGLETEKCLHVVTEAVT 61 121 PLGIYLKARVEAGGLKELEISWGLHQIVKALSFLVNDCSLIHNNVCMAAVFVDRAGEWKL 181 GGLDYMYSAQGNGGGPPRKGIPELEQYDPPELADSSGRVVREKWSADMWRLGCLIWEVFN 241 GPLPRAAALRNPGKIPKTLAPHYCELVGANPKVRPNPARFLQNCRAPGGFMSNRFVETNL 301 FLEEIQIKEPAEKQKFFQELSKSLDAFPEDFCRHKLLPQLLTAFEFGNAGAVVLTPLFKV 361 GKFLSAEEYQQKIIPVVVKMFSSTDRAMRIRLLQQMEQFIQYLDEPTVNTQIFPHVVHGF 421 LDTNPAIREQTVKSMLLLAPKLNEANLNVELMKHFARLQAKDEQGPIRCNTTVCLGKIGS 481 YLSASTRHRVLTSAFSRATRDPFAPSRVAGVLGFAATHNLYSMNDCAQKILPVLCGLTVD 541 PEKSVRDQAFKAFRSFLSKLESVSEDPTQLEEVEKDVHAASSPGMGGAAASWAGWAVTGV 601 SSLTSKLIRSHPTTAPTETNIPQRPTPEGVPAPAPTPVPATPTTSGHWETQEEDKDTAED 661 SSTADRWDDEDWGSLEQEAESVLAQQDDWSTGGQVSRASQVSNSDHKSSKSPESDWSSWE 721 AEGSWEQGWQEPSSQEPPSDGTRLASEYNWGGPESSDKGDPFATLSARPSTQPRPDSWGE 781 DNWEGLETDSRQVKAELARKKREERRREMEAKRAERKVAKGPMKLGARKLDZ

FIGURE 2

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